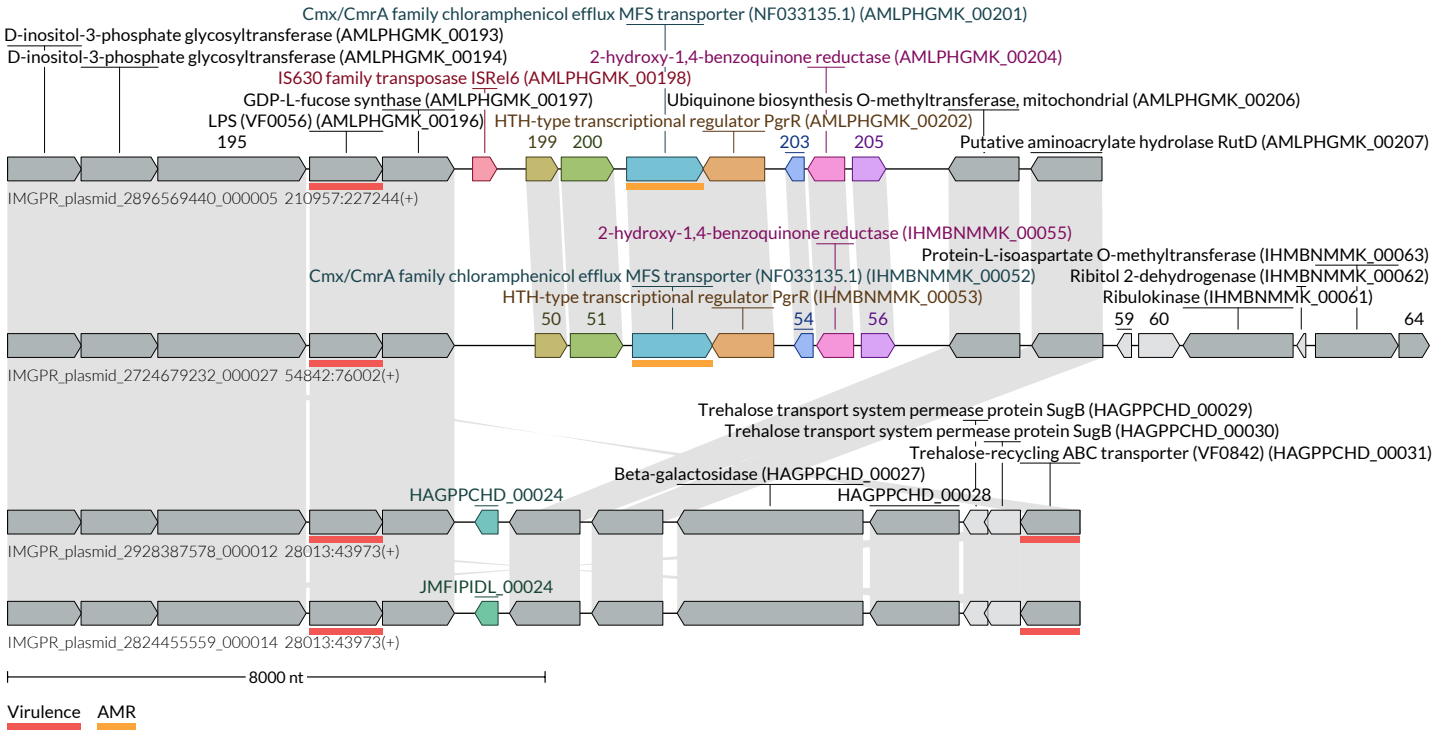
Cmx/CmrA family chloramphenicol efflux MFS transporter (NF033135.1) (AMLPHGMK_00201)
D-inositol-3-phosphate glycosyltransferase (AMLPHGMK_00193)
D-inositol-3-phosphate glycosyltransferase (AMLPHGMK_00194)
2-hydroxy-1,4-benzoquinone reductase (AMLPHGMK_00204)
IS630 family transposase ISRel6 (AMLPHGMK_00198)
GDP-L-fucose synthase (AMLPHGMK_00197)
Ubiquinone biosynthesis O-methyltransferase, mitochondrial (AMLPHGMK_00206)
LPS (VF0056) (AMLPHGMK_00196)
HTH-type transcriptional regulator PgrR (AMLPHGMK_00202)
195
199
200
203
205
Putative aminoacrylate hydrolase RutD (AMLPHGMK_00207)
IMGPR_plasmid_2896569440_000005 210957:227244(+)
2-hydroxy-1,4-benzoquinone reductase (IHMBNMMK_00055)
Protein-L-isoaspartate O-methyltransferase (IHMBNMMK_00063)
Cmx/CmrA family chloramphenicol efflux MFS transporter (NF033135.1) (IHMBNMMK_00052)
Ribitol 2-dehydrogenase (IHMBNMMK_00062)
HTH-type transcriptional regulator PgrR (IHMBNMMK_00053)
Ribulokinase (IHMBNMMK_00061)
50
51
54
56
59
60
64
IMGPR_plasmid_2724679232_000027 54842:76002(+)
Trehalose transport system permease protein SugB (HAGPPCHD_00029)
Trehalose transport system permease protein SugB (HAGPPCHD_00030)
Trehalose-recycling ABC transporter (VF0842) (HAGPPCHD_00031)
Beta-galactosidase (HAGPPCHD_00027)
HAGPPCHD_00024
HAGPPCHD_00028
IMGPR_plasmid_2928387578_000012 28013:43973(+)
JMFIPIDL_00024
IMGPR_plasmid_2824455559_000014 28013:43973(+)
8000 nt
Virulence
AMR